



[Marked up preliminary amendment]



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- 1) (Amended) A c[C]himeric expression promoter comprising at least one nucleic acid sequence, derived from a first plant promoter comprising a plant vascular expression promoter region, said plant vascular expression promoter region being replaced with a nucleic acid sequence derived from a second plant promoter and comprising a plant green tissue expression promoter region.
- 2) (Amended) The c[C]himeric expression promoter according to claim 1, wherein said first plant promoter originates from the Commelina Yellow Mottle Virus [(CoYMV)] and said second plant promoter originates from the Cassava Vein Mosaic Virus [(CsVMV)].
- 3) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1, wherein [the] said nucleic acid sequences originate from the intergenic regions of said first and second promoters.
- 4) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1, wherein it comprises at least a part of a nucleic acid sequence with SEQ ID No.1 [identified in the sequence listing under the number SEQ.ID01] fused to at least a part of a nucleic [identified in the sequence listing under the number SEQ.ID02] acid sequence with SEQ ID No.2.
- 5) (Amended) The c[C]himeric promoter [according to anyone of the preceding claims] of claim 1, wherein the nucleic acid sequence of [the] said chimeric promoter consists of a sequence selected from the group consisting of the sequences with SEQ ID Nos. 3-7 and 19-25. [identified in the sequence listing under the numbers SEQ.ID03, SEQ.ID04, SEQ.ID05, SEQ.ID06, SEQ.ID07, SEQ.ID19, SEQ.ID20, SEQ.ID21, SEQ.ID22, SEQ.ID23, SEQ.ID24 and SEQ.ID25.]
- 6) (Amended) A c[C]himeric expression promoter comprising a promoter of viral origin, of which a part consists of an exogenous element [capable of promoting] which promotes expression in plant green tissues [(GT)].

- 7) (Amended) The c[C]himeric expression promoter [according to] of claim 6, wherein [the] said [GT] exogenous promoter element is also of viral origin.
- 8) (Amended) The c[C]himeric expression promoter [according to] of claim [6] 7, wherein [the] said promoter of viral origin originates from the Commelina Yellow Mottle Virus [(CoYMV)].
- 9) (Amended) The c[C]himeric expression promoter [according to] of claim [8] 7, wherein [the] said exogenous promoter element originates from the Cassava Vein Mosaic Virus [(CsVMV)].
- 10) (Amended) The c[C]himeric expression promoter [according to] of claim 6, wherein the exogenous [GT] element replaces an endogenous vascular tissue expression [(VT)] promoter of viral origin.
- 11) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “endosperm like” box.
- 12) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “as1 like” box[ed] operably linked to [the] a plant green tissue expression GT promoter element.
- 13) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “as1” box operably linked to [the] a green tissue expression GT promoter element.
- 14) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “as2” box operably linked to [the] a plant green tissue expression GT promoter element.
- 15) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein [the one or more of the] said “as1 like”, “as1”, and “as2” boxes are operably linked upstream or downstream of the plant green tissue expression GT promoter element.

16) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein [the one or more of the] said "as1 like", "as1", and "as2" boxes are operably linked in [normal (5'>3')] 5'>3' or [inverse (3'>5')] 3'>5' orientation.

17) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it comprises at least one "as2/as2/as2" box in [normal (5'>3')] 5'>3' or [inverse (3'>5')] 3'>5' orientation.

18) (Amended) The c[C]himeric promoter [according to any one of claims 6 to 17] of claim 6, wherein it comprises at least [a] one sequence selected from the group consisting of [the] sequences with SEQ ID Nos. 3-7 and 19-25. [identified in the sequence listing under the numbers SEQ.ID03, SEQ.ID04, SEQ.ID05, SEQ.ID06, SEQ.ID07, SEQ.ID19, SEQ.ID20, SEQ.ID21, SEQ.ID22, SEQ.ID23, SEQ.ID24 and SEQ.ID25.]

19) (Amended) An e[E]xpression cassette comprising at least one nucleic acid sequence derived from a first plant promoter comprising a plant vascular expression promoter region, said plant vascular expression promoter region being replaced with a nucleic acid sequence derived from a second plant promoter and comprising a plant green tissue expression promoter region, [the sequences] said at least one sequence being operably linked to a nucleic acid sequence or gene coding for a polypeptide to be produced, said nucleic acid sequence or gene itself operably linked to a transcription termination nucleic acid sequence.

20) (Amended) The e[E]xpression cassette [according to] of claim 19, wherein said first plant promoter originates from the Commelina Yellow Mottle Virus [(CoYMV)] and said second plant promoter originates from the Cassava Vein Mosaic Virus [(CsVMV)].

21) (Amended) The e[E]xpression cassette [according to] of claim 19, wherein it comprises at least a part of a nucleic acid sequence with SEQ ID No. 1 [identified in the sequence listing under the number SEQ.ID01] fused to at least a part of a nucleic acid sequence with SEQ ID No. 2. [identified in the sequence listing under the number SEQ. ID 02.]

22) (Amended) The e[E]xpression cassette [according to] of claim 19, wherein [the] said first or second promoter [nucleic acid sequence consists of] comprises a sequence selected from the group consisting of the sequences with SEQ ID Nos. 3-7 and 19-25. [identified in the sequence

listing under the numbers SEQ.ID03, SEQ.ID04, SEQ.ID05, SEQ.ID06, SEQ.ID07, SEQ.ID19, SEQ.ID20, SEQ.ID21, SEQ.ID22, SEQ.ID23, SEQ.ID24 and SEQ.ID25.]

- 26) (Canceled) Directional desoxynucleotide building block [for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 09.
- 27) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 10.
- 28) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 11.
- 29) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 12.
- 30) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 13.
- 31) (Canceled) Directional de[s]oxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 14.

- 32) (Amended) A g[G]uide de[s]oxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to claim 1, 6, or 49 , wherein said sequence is selected from the group consisting of SEQ ID Nos. 15-18. [any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 15.]
- 33) (Canceled) Guide desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID No. 16.
- 34) (Canceled)Guide desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 17.
- 35) (Canceled) Guide desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 18.
- 36) (Amended) A v[V]ector comprising a promoter, or a promoter nucleic acid sequence, which [capable of initiating] initiates transcription of a nucleic acid sequence[, or gene,] coding for a polypeptide [to produce], wherein said[the] promoter or said[the] isolated promoter nucleic acid sequence comprises [corresponds to] a chimeric expression promoter or a promoter nucleic acid sequence according to claim 1, 6, or 49. [any one of claims 1 to 18 or 23 or 24.]
- 37) (Amended) The v[V]ector of[according to] claim 36, wherein said vector[it] is selected from the group consisting of the binary vectors pMRT1152, pMRT1171, pMRT1172, pMRT1185, pMRT1186, pMRT1187, pMRT1188, pMRT1182, pMRT1245, pMRT1246, pMRT1247, pMRT1248, pMRT1249, pMRT1250, pMRT1251, pMRT1252, pMRT1253 and pMRT1254.

38) (Amended) A t[T]ransgenic plant comprising [having] stably integrated into its genome at least one promoter or at least one promoter nucleic acid sequence according to claim 1, 6, or 49. [any one of claims 1 to 18 or 23 or 24 respectively.]

39) (Amended) The t[T]ransgenic plant of [according to ]claim 38, wherein said plant[it] is one selected from dicotyledonous species comprising[, preferably] potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, and[or] sunflower, or from monocotyledonous species comprising[, preferably] wheat, barley, oat, rice, and[or] corn.

40) (Amended) A p[P]ropagule of a transgenic plant according to claim [any one of claims] 38 or 39.

41) (Amended) The t[T]ransgenic plant propagule of[according to] claim 40, wherein it is a seed.

42) (Amended) A c[C]ell containing a promoter or a promoter nucleic acid sequence of claim 1, 6, or 49 [according to any one o claims 1 to 18 or 23 or 24 respectively], wherein said cell is selected from the group consisting of[and is preferably] a plant cell, human cell, animal cell, insect cell, bacterial cell, yeast cell, fungal cell, algal cell, and microalgal cell.

43) (Amended) The c[C]ell of[according to] claim 42, wherein it is a plant cell.

44) (Amended) A m[M]ethod for expressing a nucleic acid sequence[, or gene,] coding for a polypeptide [to produce,] by a cell, wherein said[the] method comprises[ the steps consisting of]:

- transforming said[the] cell with a vector comprising at least one promoter or at least one promoter nucleic acid sequence [according to any one of claims 1 to 18 or 23 or 24]of claim 1, 6, or 49;

- culturing said [the] cell and expressing said polypeptide encoded by said sequence in said cell [under conditions enabling the expression of the nucleic acid sequence, or gene, coding for the polypeptide].

45) (Amended) The m[M]ethod of[according to] claim 44, wherein said[the] cell is a prokaryotic or an eukaryotic cell.

46) (Amended) The m[M]ethod of according to any one of claims] claim 44 or 45, wherein said[the] cell is a cell selected from the group consisting of bacterial cells, fungal cells, yeast cells, insect cells, human cells, animal cells, algal cells, microalgal cells and plant cells.

47) (Amended) The m[M]ethod of according to any one of claims 44 to 46] claim 46, wherein said[the] cell is a plant cell.

48) (Amended) A m[M]ethod for [the] manufatur[e]ing[ of] a transgenic plant of claim 38[according to any one of claims 38 or 39], or [of] a propagule of[according to] claim 40, wherein said[the] method comprises[ the steps consisting of]:

- transforming a plant cell with a vector comprising at least one promoter or at least one promoter nucleic acid sequence of[according to any one of claims 1 to 18 or 23 or 24] claim 1, 6, or 49;

- selecting said[the] plant cell comprising[having] integrated said[the] promoter or said[the] promoter nucleic acid sequence;

- propagating said[the transformed] selected plant cell[, either] by culture or by regeneration of whole chimeric or transgenic plants.

49) (NEW) An isolated promoter nucleic acid sequence, comprising a fusion of a first sequence and a second sequence, wherein said first sequence and said second sequence comprise at least a part of each of the sequences with SEQ ID No. 1 and SE ID No. 2, respectively.

50) (NEW) The isolated promoter nucleic acid sequence of claim 23, wherein said first or second sequence comprises a sequence selected from the group consisting of sequences with SEQ ID Nos. 3-7 and 19-25.

51) (NEW) A directional deoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to claim 1, 6, or 23, wherein said sequence is selected from the group consisting of SEQ ID Nos. 8-14.